STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source:

Date Processed by STIC:

c: 08/04/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 4.4.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
 U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street,
 Alexandria, VA 22314

Revised 01/10/06



IFW16

RAW SEQUENCE LISTING DATE: 08/04/2006
PATENT APPLICATION: US/10/512,737A TIME: 10:13:05

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\08042006\J512737A.raw

- 2 <110> APPLICANT: BioTeSys GmbH 3 Schelztorstrasse 54-56 4 D 73728 Esslingen
- 5 GERMANY
- W--> 6 <120> TITLE OF INVENTION: transport system in biological systems
- C--> 7 <140> CURRENT APPLICATION NUMBER: US/10/512,737A
- C--> 7 <141> CURRENT FILING DATE: 2004-10-27
- W--> 0 <130> FILE REFERENCE:
 - 7 <150> PRIOR APPLICATION NUMBER: A 656/2002
 - 8 <151> PRIOR FILING DATE: 2002-04-29
- W--> 9 <160> NUMBER OF SEQ ID: 15

Does Not Comply Corrected Diskette Needed

ERRORED SEQUENCES

- W--> 10 <210> SEQ ID NO: 1
 - 11 <211> LENGTH: 6
 - 12 <212> TYPE: PRT
 - 13 <213> ORGANISM: Artificial sequence
- W--> 14 <220> FEATURE:
 - 15 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized
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 - 17 Gly Arg Gly Asp Ser Pro
- E--> 18 1
 - 19 <210> SEQ ID NO: 2
 - 20 <211> LENGTH: 5
 - 21 <212> TYPE: PRT
 - 22 <213> ORGANISM: Artificial sequence
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- W--> 25 <400> SEQUENCE: 2
 - 26 Tyr Ile Glu Ser Arg
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 - 28 <210> SEQ ID NO: 3
 - 29 <211> LENGTH: 5
 - 30 <212> TYPE: PRT
 - 31 <213> ORGANISM: Artificial sequence
- W--> 32 <220> FEATURE:
 - 33 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized
- W--> 34 <400> SEQUENCE: 3
 - 35 Ala Asp Gly Glu Ala
- E--> 36 1
 - 37 <210> SEQ ID NO: 4

Invalid aid Amino aid Numbering

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RAW SEQUENCE LISTING DATE: 08/04/2006 PATENT APPLICATION: US/10/512,737A TIME: 10:13:05
                     Input Set : A:\pto.da.txt
                     Output Set: N:\CRF4\08042006\J512737A.raw
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     53 Val Arg Leu Leu Asn Asn Trp Asp
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     66 <212> TYPE: PRT
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     76 <212> TYPE: PRT
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     81 Leu Ser Gly Ala Leu Arg
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     83 <210> SEQ ID NO: 9
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     85 <212> TYPE: PRT
     86 <213> ORGANISM: Artificial sequence
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Barrel

DATE: 08/04/2006

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PATENT APPLICATION: US/10/512,737A
                                                           TIME: 10:13:06
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                    Output Set: N:\CRF4\08042006\J512737A.raw
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E--> 92 1
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                            _5 10
E--> 93 15
    94 Leu Phe Val Met Trp Met
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     99 <213> ORGANISM: Artificial sequence
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    115 Leu Phe
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RAW SEQUENCE LISTING

RAW SEQUENCE LISTING

DATE: 08/04/2006

PATENT APPLICATION: US/10/512,737A

TIME: 10:13:06

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\08042006\J512737A.raw

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139 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized

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W--> 149 <400> SEQUENCE: 15

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E--> 151 1

Source Source

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 08/04/2006 PATENT APPLICATION: US/10/512,737A

TIME: 10:13:07

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\08042006\J512737A _raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:9; Line(s) 92 Seq#:11; Line(s) 113

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/512,737A TIME: 10:13:07

DATE: 08/04/2006

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\08042006\J512737A.raw

L:6 M:283 W: Missing Blank Line separator, <120> field identifier L:7 M:270 C: Current Application Number differs, Replaced Current Application No L:7 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:0 M:201 W: Mandatory field data missing, <130> FILE REFERENCE L:9 M:283 W: Missing Blank Line separator, <160> field identifier L:10 M:283 W: Missing Blank Line separator, <210> field identifier L:14 M:283 W: Missing Blank Line separator, <220> field identifier L:16 M:283 W: Missing Blank Line separator, <400> field identifier L:18 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1 L:23 M:283 W: Missing Blank Line separator, <220> field identifier L:25 M:283 W: Missing Blank Line separator, <400> field identifier L:27 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:2 L:32 M:283 W: Missing Blank Line separator, <220> field identifier L:34 M:283 W: Missing Blank Line separator, <400> field identifier L:36 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3 L:41 M:283 W: Missing Blank Line separator, <220> field identifier L:43 M:283 W: Missing Blank Line separator, <400> field identifier L:45 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:4 L:50 M:283 W: Missing Blank Line separator, <220> field identifier ~1:52 W#263 W: Missing Blank Line separator, <400> field identifier t L:54 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:5 L:59 M:283 W: Missing Blank Line separator, <220> field identifier L:61 M:283 W: Missing Blank Line separator, <400> field identifier L:63 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6 L:68 M:283 W: Missing Blank Line separator, <220> field identifier L:70 M:283 W: Missing Blank Line separator, <400> field identifier L:72 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7 L:78 M:283 W: Missing Blank Line separator, <220> field identifier L:80 M:283 W: Missing Blank Line separator, <400> field identifier L:82 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:8 L:87 M:283 W: Missing Blank Line separator, <220> field identifier L:89 M:283 W: Missing Blank Line separator, <400> field identifier L:92 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:9 L:93 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:9 L:95 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:9 L:100 M:283 W: Missing Blank Line separator, <220> field identifier L:102 M:283 W: Missing Blank Line separator, <400> field identifier L:104 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:10 L:109 M:283 W: Missing Blank Line separator, <220> field identifier L:111 M:283 W: Missing Blank Line separator, <400> field identifier L:113 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:11 L:114 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:11 L:120 M:283 W: Missing Blank Line separator, <220> field identifier L:122 M:283 W: Missing Blank Line separator, <400> field identifier L:124 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:12 L:129 M:283 W: Missing Blank Line separator, <220> field identifier L:131 M:283 W: Missing Blank Line separator, <400> field identifier L:133 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:13

VERIFICATION SUMMARY

DATE: 08/04/2006

PATENT APPLICATION: US/10/512,737A

TIME: 10:13:07

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\08042006\J512737A.raw

L:138 M:283 W: Missing Blank Line separator, <220> field identifier L:140 M:283 W: Missing Blank Line separator, <400> field identifier L:142 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:14 L:147 M:283 W: Missing Blank Line separator, <220> field identifier L:149 M:283 W: Missing Blank Line separator, <400> field identifier L:151 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:15